

SIMULATE PROCESS FLOW

Help Document

Sim.Pro.Flow is a decision support tool that automates the build of a discrete event simulation and allows for mapping, modelling and improving of system pathways.

This help document will describe the function of all the widgets within Sim.Pro.Flow (v2.0). The document is structured where each section discusses a main tab, and the subsections discuss any subsequent tabs. This document does not provide instructions of how to use Sim.Pro.Flow but will address the considerations required by the user.

Each widget has a corresponding numbered item indicated by either a blue or purple circle, where is blue circle is standard. A purple circle indicates that a document will be produced and is described in the Output Files section.

Images outlined with a thick black dashed line indicates that this is a snippet rather than the full screen view.

The sample datasets shown were artificially generated. The dataset from Data Type 1 is used for examples shown, which consists of 16 fictional pathways that each occur once.

Note: >> indicates path, " indicates fixed name, [] indicates description of name that will change, bold text will indicate a point in the image.

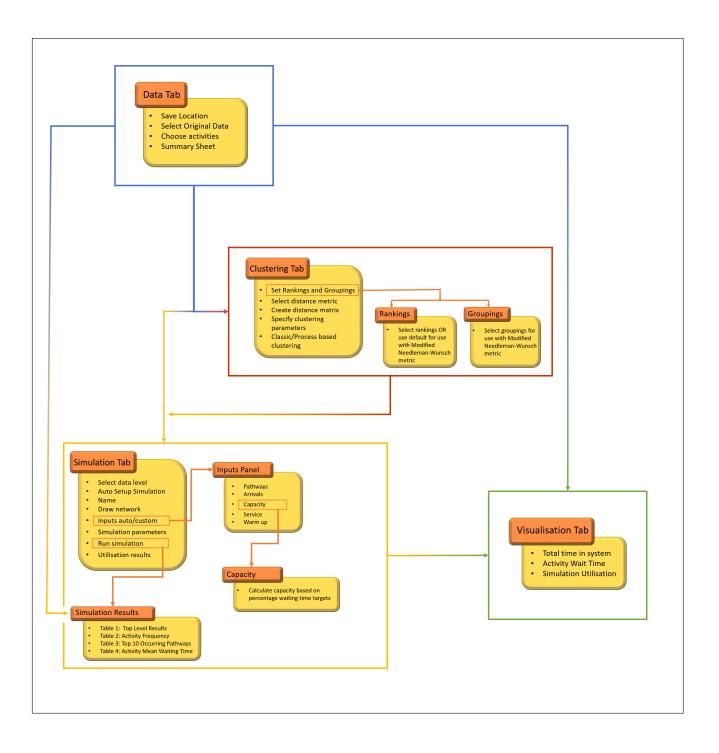
Note: Terms defined in the glossary will be linked once on each page it occurs.

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1 Usage Map

The usage map displays the four main tabs within Sim.Pro.Flow and how information moves between them.



2 Data Panel



1) File Menu

- About Opens the about box for Sim.Pro.Flow.
- Help Opens this help document.
- Export Saves the raw simulation variables.
- Save Saves the four main results tables.

2) Select Save Location

Opens the file explorer window.

Select the folder to save the information for this session. On selection multiple documents will be produced. *Suggested to create a new folder for each session.*

3) Select Data

Opens the file explorer showing Excel files (.xlsx). Select the dataset for this session.

4) Select Columns

Opens the Select Columns pop-up box (5).

5) Select Columns Pop-up Box

Allows the user to select the columns required for analysis with Data Type 1 or Data Type 2.

Select all button checks all column checkboxes.

Select the **Group Codes** checkbox to indicate Data Type 2.

Only columns containing dates should be selected. Do not select blank columns.

**A copy of the data will be saved in the save location, with naming convention SimProFlow_DataFileName. **

6) Format

To be used with Data Type 3. Three drop down boxes will be populated with all column headers, for the user to select the corresponding column for **Unique id**, **Activity** and **Date**.

A copy of the data will be saved in the save location with naming convention 'SimProFlow_'[DataName].

7) Create Summary Sheet

Produce a word document 'Summary_Sheet.docx'.

8) Table

After the column selection has taken place (either through (5) or (6)) a non-editable table will be created displaying the activity name and the corresponding code.

3 Clustering Panel



1) Set Rankings and Groupings

Populates the Rankings and Groupings subtabs for use with the Modified Needleman-Wunsch metric.

2) Select Penalty Values

Spin control to allow user to select the penalty values for use with the Modified Needleman-Wunsch metric. Has lower and upper bounds of 0 and 100 respectively.

3) Select Distance Metric

Drop down box containing the eight supported distance metrics, namely: Levenshtein, Damerau-Levenshtein, Jaro, Jaro-Winkler, Needleman-Wunsch, Jaccard, Cosine, Longest Common Subsequence, and Modified Needleman-Wunsch.

If Modified Needleman-Wunsch chosen, Rankings and Groupings must have been selected.

4) Create Matrix

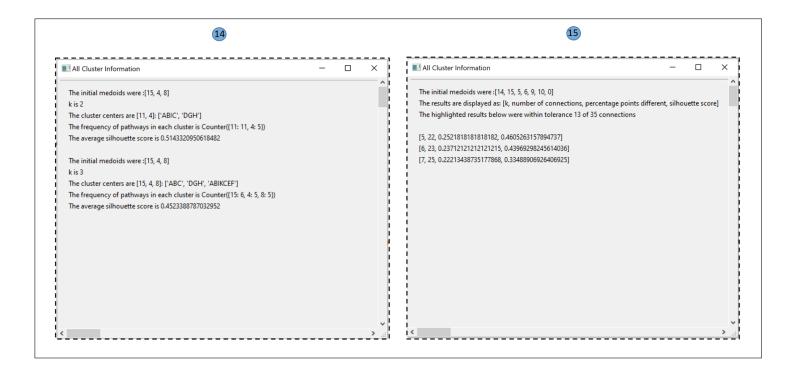
A ixi distance matrix will be created using the chosen distance metric i.e. the distance from each pathway in the dataframe to every other pathway will be calculated.

5) Cluster Centroids

There are five methods supported for choosing the k number of initial centroids, where k is as defined in (6):

- Random: Random set of unique values corresponding to pathway indexes.
- Most Occurred: Index of the pathways that occurred most often. By the definition of the dataframe, this will be the first k pathways.
- Least Distance: Pathway indexes that generated the smallest value of the sum of it's row.
- Specify: Allows the user to specify pathway indexes in the 'Specify' input box. There must be k comma separated values specified within the range [0, i].
- Previous: Select a preciously used set of initial centroids from the Choose Previous drop down box. This will be populated after each clustering is complete, thus can not be selected before the first clustering has occurred. The selected set must contain the same number of values as the specified k.

. **Only one method can be selected, otherwise an error box will appear. If multiple selected post error box then the topmost selected method will be used. **



6) Cluster Results

Spin control box to select max k within range [2, 100]. Select the level of highlighted results to be displayed.

- All: Display all results in the range [2, max k]. This selection will not be able to be used with subsequent simulation.
- Besk k: Display the results for the k producing the best Silhouette Score in the range [2, max k].
- Besk k: Same as previous but k=2 can not be selected.
- k only: Display the results for the value of k specified in the max k spin control.

7) Select Process Based Percentage

For comparing clustering results transition matrix with the adjusted transition matrix.

Selected **Adjust percentage** will be used for the adjusted transition matrix. *Used in process based clustering only.*

8) Select Process Based Tolerance

Used to specify which results to display in (15).

9) Save Centroids

Saves the cluster centroids to the Excel file.

Results saved will correspond to the results choice selected in (6).

10) Classic Cluster (14)

k-medoids clustering applied to the distance matrix using the specifications selected in points (5) - (9). The resulting k will be used as the number of transition matrices to be used with the Clustered Transitions simulation.

11) Process Based Cluster (15)

k-medoids clustering applied to the distance matrix using the specifications selected in points (5) - (9). The resulting k will be used as the number of medoids to be used with the Process Medoids simulation.

12) Canvas

Operates as universal widget.

Displays process based violin plot to aid k selection.

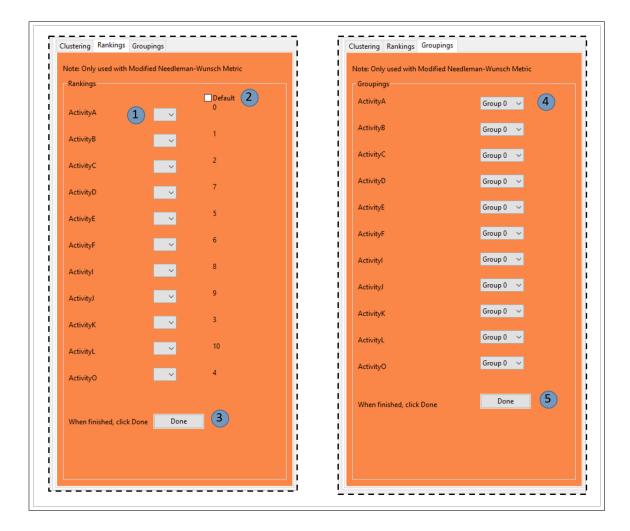
Naming convention:

[Metric]_[InitialCentroidSet]_[ResultsType]_[k]

13) Toolbar

Operates as universal widget.

3.1 Rankings and Groupings



1) Select Rankings

Rankings to be used with Modified Needleman-Wunsch metric.

Drop down box to select rank for activity, where options available are [0, N].

Lower value corresponds to more important.

Choose unique value for each activity.

2) Use Default

Check box to use default rankings shown.

Default rankings calculated based on frequency of activity.

3) Confirm Rankings

Done button to confirm rankings.

This must be pressed to use Modified Needleman-Wunsch metric.

4) Select Groupings

Groupings to be used with Modified Needleman-Wunsch metric.

Drop down box to select group for activity, where options available are [Group 0, Group N].

Only activities in the same group will be able to be swapped by definition of the Modified Needleman-Wunsch metric.

5) Confirm Groupings

Done button to confirm groupings.

This must be pressed to use Modified Needleman-Wunsch metric.

4 Simulation Panel

4.1 Model



1) Select Data Level

Select the data level to use for the simulation. Drop down options: Raw Pathways, Full Transitions, Clustered Transitions and Process Centroids.

2) Auto Setup Simulation

Construct the simulation for selected data level.

Must be initially pressed to start using this tab.

Should be pressed to change data level as selected.

For clustered transitions and process medoids will use the most recent results. Please ensure that this was not the results type 'All'.

3) Provide Simulation Name

Name for simulation. Automatically filled on **Auto Setup Simulation** based on data level with convention [Initial]_Basic, where Initial = R, F, C, P respectively. *Must be unique for each execution of simulation - prompted by error.*

**If trials > 1 then name appends $_Trials_[No.Runs]$ **

4) Network: Draw

Draw: Draw the networks for the selected data level. Drop down box: Contains names of networks produced. View: Open the selected network pdf in a pdf viewer.

5) Right Hand Panel

After auto setup simulation panel will populate as shown in Custom Inputs.

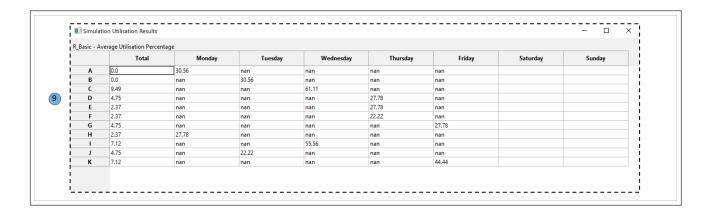
6) Simulation Custom Inputs

Use auto or custom values for simulation inputs.

7) Time Elements

Simulation parameters relating to time.

- Week Type: Number of working days in a week. Options 5 days, 7 days. Used in Capacity simulation input.
- Number of Individuals: Simulate until number of individuals specified have completed simulation.
 Auto - Number of individuals in data,
 Custom - defined in Arrivals.
- Target Days: Target days for time in system. Used in tabs Capacity and Simulation Results.
- Number of Trials: Number of simulation runs. *Simulation seed set to the corresponding run number starting 0.*
- Simulation Seed: Seed used for the simulation. *Not used if Number of Trials > 1.



8) Run Simulation

Runs the simulation.

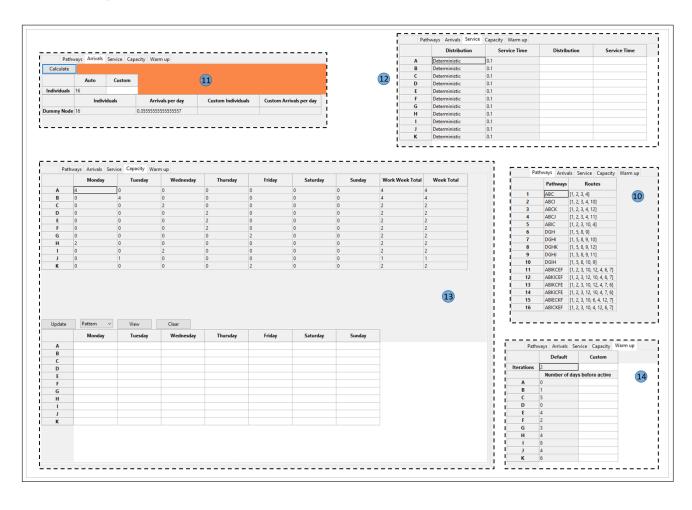
Pop up box will confirm when simulation is complete.

9) View Utilisation Results

Opens Simulation Utilisation Results pop up box. Displays the mean average percentage of capacity used on each named day.

Not produced for trials.

4.1.1 Custom Inputs



From the simulation model tab the simulation inputs (6) can be selected as either auto or custom. Both the auto and custom options are hosted within this right hand panel (5) and initially appear after auto setup simulation (2) has been performed. The pathways and arrivals are data level specific and shall be discussed for each, whereas the remaining inputs are universal across all data levels.

Note: The example shown are for raw pathways. Note: Any cell in grey is non editable.

1) Pathways

This displays the pathways that will be taken by the individuals.

- Raw Pathways: Table will contain two columns Pathways and Routes.
 - The pathways will correspond with the dataframe and the routes will reflect the route to take through the nodes of the pathways.
 - The raw pathways all enter the simulation through a dummy node (node 1) and thus all pathways will start with 1, with the resulting nodes corresponding to letter code position +1.
- Full Transitions: This will displays a N xN transition probability matrix.
- Clustered Transitions: Takes the results from the classic clustering for k clusters and produces a NxN transition probability matrix for each. Each cluster will be represented by a class.
- Process Medoids: Takes the results from the process based clustering for k medoids and displays the table as is for the Raw Pathways, where the pathways correspond to the k medoids.

2) Arrivals and Number of Individuals

The arrivals correspond to the arrival rate for the simulation. The value/s for the arrivals per day will be used as the λ for an Exponential distribution. The general arrival rate is defined in Equation 1. This is only editable through the number of individuals, where the user can enter the number of individuals in the 'Custom' cell and on execution of the 'Calculate' button the arrival rate will be re-calculated. The user can then choose to either simulate until the auto or custom individuals and use the auto or custom arrivals independently.

- Raw Pathways: Arrivals will be received at a dummy node (node 1) where the individual will be assigned their pathway.
 - Arrival equation applied where $A_n = N$.
 - *The individuals pathway number will correspond to the individuals id number.*
- Full Transitions: Individuals will arrive into respective arrival activities as extracted from the transition probability matrix.
 - Table will have N rows, one for each activity, with arrival equation calculated for each.

- Clustered Transitions: Same as the Full transitions with the addition that the columns will be repeated k times i.e. arrival matrix for each class. Arrival equation values for A_n are obtained from the propagated values.
- Process Medoids: Table will have k rows where the arrival activity will correspond to the first activity of the medoid for each class.

 Arrival equation values for A_n are obtained from the propagated values.

3) Service

Service rates are auto set to deterministic at 0.1. Custom distributions can be deterministic or exponential. Service time can take a float value and will be formatted as necessary for the chosen distribution.

4) Capacity

Top table displays the auto calculated capacity for each activity - the average number of individuals seen on each named day .

Bottom table is used for capacity where users can manually input custom values for capacity.

Various custom options are available to auto-fill the custom table using the following widgets:

- Update: Update options in the drop down box.
- View: Populate bottom table with chosen option.
- Clear: Clear the table.

Three custom option types are available which will consider the week type (5 days or 7 days) chosen:

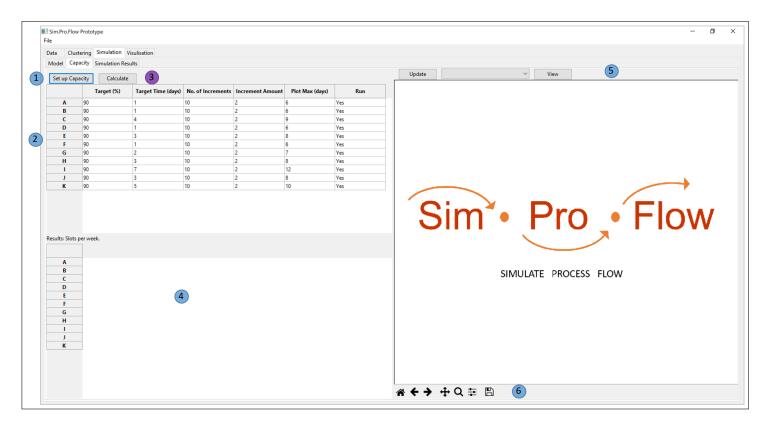
- Pattern Use the auto pattern from the top table.
- Smoothed Spread the total week capacity (considering chosen week type) evenly across the working days.
- Naming Convention: Cal_Cap_[Number] Takes the results calculated in the Capacity tab and uses the smoothed technique.

5) Warm-up

Three warm up types available:

- None: No warm up used. This is the default.
- Iterative: Collect results once w multiples of I have completed the simulation. Auto set to 1.
- Warm Start: Block service at activity for an initial number of days. Auto set to original mean activity waiting time (Table 4).

4.2 Capacity



This calculation, developed by Arruda et al [1], will suggest the amount of weekly capacity for each activity required to allow a percentage of individuals to wait no more than a target number of days e.g. 90% of individuals to wait no more than 10 days. Various values of capacity will be tested, based on the inputs provided, to suggest the number of slots per week required. The base value is auto calculated from the activity demand.

1) Set up Capacity

Create and populate capacity Inputs Table (2).

2) Inputs Table

Table to control inputs for capacity calculation.

- Target (%): Percentage to wait no longer than **Target Time**.
- Target Time (days): The target number of waiting days.

Set up Capacity will auto populate this with the floor of the average waiting time for each activity in the original data i.e. Table 4.

The minimum value allowed is 1.

- No. of Increments: This will set the number of values to explore.
- Increment Amount: This will set the amount to increment the base value by.
- Plot Max (days): This is the maximum number of days to view on the plot.
 Auto populated to Target Time + 5.

• Run: Choose if Yes/No to run the calculation for this activity.

3) Calculate

This performs the calculation described in [1].

4) Results

A value indicating the number of capacity slots required each week for each activity will be produced.

*If No selected for an activity, the previous value will be shown in the **results table (4)** in red font.*

If No selected for first run then red 0 will appear in results table.

5) Canvas

Operates as universal widget.

Displays line chart showing the percentage waiting no more than x (axis value) days.

Naming convention: Cal_Cap_NumberOfCalculations

6) Toolbar

Operates as universal widget.

4.3 Simulation Results



1) Update Tables

Update button will update all tables with any new results that have been produced since the last update. On update Table 1 will append rows, all other tables will append columns.

The first entry for all the tables will correspond to the original input data and is named original

2) Table 1

Summary Results:

- Mean Time in System: Mean average time from arrival to exit in days.
- Median Time in System: Median average time from arrival to exit in days.
- Target [days, %]: Percentage total time in system less than the specified target days.
- No. Unique Pathways: Number of unique pathways performed.
- Occurs Once: Number of pathways that occur only once.
- Occurs > Once: Number of pathways that occur more than once.
- Total Transitions: The total sum of the difference matrix.

- Mean Transitions: The mean average of the difference matrix.
- Largest Transitions: The largest entry in the difference matrix.
- Day Last Arrival: The day that the last arrival occurred.
- Overall Period: The total number of days that occurred.
 - *For the original data this is calculated by latest date recorded earliest date recorded.*

3) Table 2

Activity Frequency: The number of times each activity was performed.

4) Table 3

The top 10 occurring pathways and their corresponding frequency count.

Only pathways occurring more than once will appear.

5) Table 4

Activity Wait times (days): Mean average wait time for each activity in days.

5 Visualisation Panel



1) Total Time Tab

Displays a of histogram for total time in system, where the left canvas is for the original data and the right canvas is for the selected simulation.

The original data canvas is possible to view before any simulation has been run

2) Waiting Time Tab

Displays a plot containing multiple histograms displaying the waiting time for each activity, where the left canvas is for the original data and the right canvas is for the selected simulation.

The original data canvas is possible to view before any simulation has been run

3) Utilisation Tab

Displays two line charts of the capacity utilisation, where the left canvas plots the percentage of the capacity utilised each day and the right canvas plots the number of individuals remaining in the queue at the end of each day.

12) Canvas

Operates as universal widget.

Drop down options will be populated with simulation names.

13) Toolbar

Operates as universal widget.

Note: *After view pressed, each canvas will required manual updating by selecting the move tool and clicking in the canvas area*

Note: **The plot axis scale and subplot configuration is automated to best suit each plot.**

Note: ***These plots are not available for trial simulations. ***

6 Auxiliary Information

6.1 Input Data

		ActivityA	ActivityB	ActivityC	ActivityD	ActivityE	ActivityF	Activityl	ActivityJ	ActivityK	ActivityL	ActivityO		
			02/01/2020											
			02/01/2020							10/01/2020	1			
			02/01/2020									12/01/2020		
	1		02/01/2020								09/01/2020			
		01/01/2020	02/01/2020	17/01/2020						10/01/2020)			
					04/01/2020			05/01/2020	08/01/2020					
					04/01/2020			05/01/2020	08/01/2020	10/01/2020)			
					04/01/2020			12/01/2020	15/01/2020			19/01/2020		
					11/01/2020			12/01/2020	15/01/2020		16/01/2020)		
					11/01/2020				22/01/2020					
		08/01/2020	09/01/2020	24/01/2020			25/01/202			17/01/2020		19/01/2020		
			09/01/2020				25/01/202			17/01/2020		12/01/2020		
			09/01/2020			08/02/2020				24/01/2020		26/01/2020		
			09/01/2020				08/02/202			31/01/2020		26/01/2020		
			09/01/2020				08/02/202			24/01/2020		02/02/2020		
							08/02/202			24/01/2020		02/02/2020		
		115/01/2020												
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Type 1

1) Original Input Data

Each row corresponds to an individual. Each activity has it's own column.

Date stamps to record activity and left blank otherwise.

2) SimProFlow_[DataName] - Data Tab

Additional information appended to **Original Input Data** and saved in the save location as an Excel file with naming convention SimProFlow_[DataName]. Description of the columns:

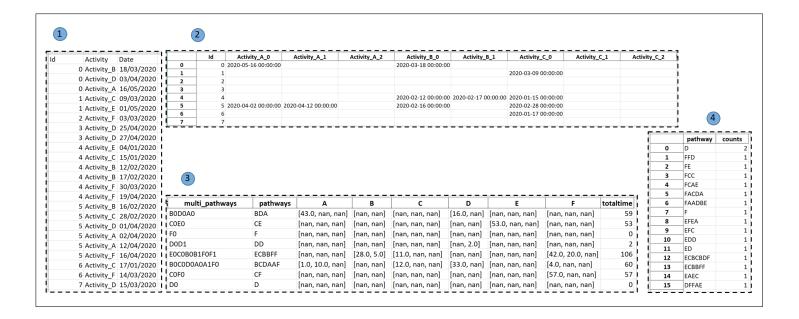
- pathways: Displays the string representing the pathway taken by the individual.
- [Letter]: Waiting time for performed activity, in days.
- totaltime: This is the total waiting time in the system, in days.

3) $SimProFlow_[DataName]$ -Dataframe Tab

This contains the unique pathways and the number of times they were performed (counts).

This will be sorted by decreasing count then alphabetically decreasing.

^{*}Do not select any blank columns.*



Type 2

1) Original Input Data

Three columns:

- Id: Containing unique identifier for an individual. *This can be alphanumeric.*
- Activity: Containing activity names.
- Date: Containing performance date.

Columns do not have to have these specific titles.

**Dataset can contain additional (redundant)

columns.**

2) SimProFlow_[DataName] - Data Tab

Converted **Original Input Data** into the format where each row corresponds to an individual and each column corresponds to a performance of an activity. [Activity name] followed by [_Number] corresponds to the performance occurrence number, starting at 0 e.g. _0 is the first time, _1 is the second time.

This should correctly correlate to lower [_Number] and earlier date.

3) SimProFlow_[DataName] - Data Tab Cont. Additional information appended to Original Input Data and saved in the save location as an Excel file with naming convention SimProFlow_[DataName]. Description of the columns:

- multi_pathways: This will contain the pathway string consisting of the letter code followed by the [_Number].
- pathways: Displays the string representing the pathway taken by the individual.
- [Letter]: Waiting time for performed activity, in days.

This will be a list of values the length of the maximum number of times one individual performed that activity e.g. Column D contains lists of length 2 as pathway 3 (D0D1) contains two performances of activity D.

nan corresponds to either the activity wasn't performed or it was the first activity and doesn't have a waiting time e.g. pathway 1 (C0E0) has list [nan, nan, nan] for C even though C was performed, as this was the first activity.

• totaltime: This is the total waiting time in the system, in days.

4) SimProFlow_[DataName] - Dataframe Tab

This contains the unique pathways and the number of times they were performed (counts).

This will be sorted by decreasing count then alphabetically decreasing.

Type 3

This will take in data of the form (2) and group together activities with the same name but differing [Number]. For example Activity_A_0, Activity_A_1 and Activity_A_2 will be considered the same activity.

6.2 Universal Widgets

Two widgets are repeatedly used throughout the interface.

Canvas

The panel is made up of:

- Update button: Update the options available in the drop down box.
- Drop Down box: Show the options available to be viewed.
- View: Load the selected option into the canvas.

 *The option will not appear on the canvas until it
 has been activated via a Toolbar item.*

All options available to view will be saved in a corresponding location within the save location and called upon to view.

The Sim.Pro.Flow logo is used as a placeholder.

Toolbar

This is matplotlib's interactive navigation toolbar [12]. The two main tools allow the user to move around the canvas and zoom into a selected area.

To move: Select the cross arrow to click and drag around the image.

This item can be used to activate the canvas.

To zoom: Select the magnifying glass to click and drag to form a box which will enlarge to fill the canvas.

The home tool will reset the image and the back and forward arrows will move between the previous and next actions performed respectively.

It is not necessary to use the save tool as all images will have been saved in the save location.

6.3 Data Levels

There are four possible data levels to be used within Sim.Pro.Flow. Each data level explores a deeper granular level for mapping the pathways. Each data level is described in turn, including the advantages and disadvantages, to help the user decide the most appropriate data level to use.

- Raw Pathways: These are the pathways exactly as they appear in the data. The purpose of using the raw pathways is to allow validation of the simulation parameters, by ensuring that the key performance indicators (KPI's) are reflective of those seen in the data. This is a major advantage and eases the user in to a system exploring pathways that are familiar and have actually previously happened in reality. The main disadvantage of this level is that it is very specific to the input data and doesn't allow for any variation.
- Full Transitions: The transition matrix can be extracted from the raw pathways. This allows for some variation to be introduced into system, which can be more reflective of future events. The disadvantages are that this can lead to unrealistic and impossible pathways. As the movements through the system will be based off probabilities extracted from the data, it could lead to individuals bouncing around within the system and performing really long pathways. Furthermore, restricting an activity to be performed at most once (referred to as at most once restriction) can not be enforced here.
- Cluster Transitions: From the results of the clustering the transition matrix for each cluster can be extracted. The aim for this level is to have more precise transition matrices, as the pathways have been clustered with those that they are most similar to, and thus resulting pathways should display less variation than the full transitions. This method has all the same disadvantages as the Full Transitions.
- **Process Medoids:** The second piece of usable information that is produced from the clustering are the pathways that are selected as the medoids. These exact pathways can be performed, and as such solves the disadvantages of using the transition matrices, as now only previous pathways can be performed and if the input data has the at most once restriction, so will the process medoids. Similar to the Raw Pathways, this does pose the lack of variation disadvantage but now with even more restriction.

6.4 Process Based

The process based clustering will perform the same process as the class clustering (using k-medoids), however the results of the clustering will be reported and used differently. The process based will use only the pathways selected as the medoids for each cluster. Therefore extra information is provided to support the user to select the number of k clusters to use.

This will report values to represent how closely the medoids reflect the original pathways. Putting this in terms of the network, this would translate to minimise the number of edges whilst also minimising the differences in the edge values in comparison to the full network.

Selection

The calculation process is as follows:

- 1. In the original data matrix: Remove transition occurring less than a specified number of times,
- 2. Calculate this adjusted transition matrix,
- 3. From the medoids, construct the transition matrix, accounting for connections representing the number of pathways in the cluster,
- 4. Calculate the absolute difference matrix,
- 5. Count the number of non-zero connections,
- 6. Calculate the average percentage points different,
- 7. Plot all non-zero differences.

Output Files 6.5

Note: >> indicates path, " indicates fixed name, [] indicates description of name that will change. Note: Main location is the selected save location folder.

Action Button	Name	Description
Housin Barron	Data Tab	Description
	Folder 'Network_diagrams'	Empty folder for Draw to output.
	Folder 'Plots	General folder to contain place specific folders for
		plots.
	'Plots'	Empty folder to contain plots from the Capacity
	>> 'Capacity' Folder	tab.
	'Plots'	Empty folder to contain the process violin plots
	>> 'Process_Violin_Plots' Folder	from the clustering tab.
	'Plots'	Empty folder to contain all plots from the simu-
	>> 'Simulation' Folder	lation tab.
	'Plots'	Empty folder to contain plots from running trials
Select Save	>> 'Trials' Folder	in the simulation tab.
Location	'Plots'	Empty folder to contain the plots for the sum-
	>> 'Summary' Folder	mary sheet.
	'Clustering_Transition_Matrix.xlsx'	Empty excel file to add sheets for the clustering
	'Process_Centroids.xlsx'	tansition matrix from the clustering tab.
	Process_Centroids.xisx	Empty excel file to add sheets for the process
	'Raw_Sim_Results.xlsx'	based clustering centroids from the clustering tab. Empty excel file to add sheets for the raw simu-
	rtaw_Sim_rtesuits.xisx	lation results from the simulation tab.
	'Simulation_Difference_Matrix.xlsx'	Empty excel file to add sheets for the simulation
	Simulation_Difference_Matrix.xisx	difference matrix from the simulation tab.
	'Cluster_Centroids.xlsx'	Empty excel file to add sheets for the classic clus-
	0-100 00-100 00-100	tering centroids from the clustering tab.
	'SimProFlow_[DataName].xlsx'	Additional information added to a copy of the
		original data file, as shown in data types.
	'Plots' >> 'Simulation'	Plot of subplots containing histogram of the wait-
Select Columns/	>> 'Activty_Waits_original.png'	ing time for each activity in the oringal data se-
Format		lection.
	'Plots' >> 'Simulation'	Histogram of the total time in system for the orig-
	>> 'TotalTime_original.png'	inal data.
	'Summary_Sheet.docx'	Word document produced containing summary
		information about the original data. Some of the
		information included is the number of individuals,
		number of pathways, mean, median and quartile
	171	time in system, as well as the following four plots.
G	'Plots' >> 'Summary'	Horizontal bar chart displaying the number of
Create Summary	>> 'Activity_Frequency.png'	times (frequency) that each activity was per-
Sheet	'Plots' >> 'Summary'	formed. Revealet of the activity waiting times
	>> 'Boxplot_Activity_Wait_Time.png'	Boxplot of the activity waiting times.
	'Plots' >> 'Summary'	Heatmap displaying a representation of all path-
	>> 'Heatmap_All_Pathways.png'	ways from the original data.
	'Plots'>> 'Summary'	Histogram of the total time in system. *This may
	>> 'Histogram_Total_Time_in_System.png'	be a different scale to 'TotalTime_original'.*
	// moogram_rotal_rme_m_bystem.pmg	or a angerena scare to 10th 1 the of tymul.

Action Button	Name	Description						
Clustering Tab								
Create Matrix	Update 'Clustering_Transition_Matrix.xlsx'	Sheet added with distance metric code as sheet name. If Modified Needleman-Wunsch selected sheet name is MNW_[mgsns] where m, g, s and ns are the penalty values selected. Contains nxn distance matrix for pathways in same order as dataframe.						
Save Centroids	Classic Cluster Used: Update 'Cluster_Centroids.xlsx'	On first save create sheet Set_Medoids recording the set number and the initial centroids. This will subsequently be updated. Sheet added called [Metric]_Set_[SetNumber]_df containing all the pathways in the dataframe and the index of the corresponding medoids of the cluster it belongs to.						
Checked	Process Cluster Used: Update 'Process_Centroids.xlsx'	On first save create sheet Set_Medoids recording the set number and the initial centroids. This will subsequently be updated. Sheet added called [Metric]_Set_[SetNumber] will contain the medoids for k (based on results type) and the number of pathways assigned to each medoids. Sheet added called [Metric]_Set_[SetNumber]_df containing all the pathways in the dataframe and the index of the corresponding medoids of the cluster it belongs to.						
Process Cluster	'Plots' >> 'Process_Violin_Plots' >> [Metric_Set_SetNumber_Type_k].png	Creates the violin plot for all values in [2, max k] to aid in decision of k values to use.						

Simulation Tab

	Update 'Raw_Sim_Results.xlsx'	Adds a sheet called [SimName] containing the raw simulation results. Each row is an individual with columns for id, waiting time at each activity, pathway, total time in system and customer class.
Run Simulation	Update 'Simulation_Difference_Matrix.xlsx'	Initially adds sheet called original. For each simulation adds a sheet called [Sim-Name] containing the transition matrix excluding the start transitions and including the end transitions.
Tun Simulation	'Plots' >> 'Simulation' >> 'Activity_Waits_'[SimName].png 'Plots' >> 'Simulation'	Plot contianing subplots of histograms for the waiting time for each activity. Histogram of the total time in system.
	>> 'TotalTime_'[SimName].png	
	'Plots' >> 'Simulation' >> 'Utilisation_Percent_'[SimName].png	Plot containing subplots of a line chart showing the percentage of capacity used each day for each activity.
	'Plots' >> 'Simulation' >> 'Utilisation_Queue_'[SimName].png	Plot containing subplots of a line chart showing the number of individuals remaining in the queue at the end of each day.

Action Button	Name	Description
	*Note in all draw diagrams a light grey line with	_
	whether that be one connection or a	a probability of 1.*
Auto Setup - Full Transitions - Draw	'Network_diagrams' >> 'Network_'[SimName].png	Directed graph of the full transitions network. Each node is an activity where an edge between two nodes represents the transition probability. The start and end nodes are included for visual aid. The start node itself is not the arrival node for the simulation.
Auto Setup - Clustered Transitions - Draw	'Network_diagrams' >> 'Network_'[SimName]'_Class'[_No.].png	Directed graph for each class for the clustered transitions. There will be a file created for each class/cluster where [No.] is the class number. Each node is an activity where an edge between two nodes represents the transition probability. The start and end nodes are included for visual aid. The start node itself is not the arrival node for the simulation.
	'Network_diagrams' >> 'Network_'[SimName_k_].png	Directed graph for the process medoids. Each node is an activity where an edge between two nodes represents the transition probability from within the set of medoids. The start and end nodes are included for visual aid. The start node itself is not the arrival node for the simulation.
Auto Setup - Process Based - Draw	'Network_diagrams' >> 'Network_'[SimName_k_]'_adjust'[_Perc].png	two nodes represents the transition probability for the original transitions after adjusting for the adjust percentage ([Perc]). The start and end nodes are included for visual aid. The start node itself is not the arrival node for the simulation.
	'Network_diagrams' >> 'Network_'[SimName_k_]'_pathways'.png 'Network_diagrams' >> 'Network_'[SimName_k_]'_linked'.png	Visual representation of the raw medoids. Visual representation of the raw medoids where activities are grouped by position. The boxes define the position of the activity (reading left to right). The boxes contain nodes representing all activities that occurred at that position in the set of medoids. The connections between activities in the groups represent the number of times that connection occurred in the set of medoids.
	Simulation Conscitu	Tab
Calculate	Simulation - Capacity 'Plots' >> 'Capacity' >> 'Cal_Cap_'[Number].png	Plot containing subplots for each activity displaying the percentage seen within x axis days for the various values of weekly capacity investigated.
	File Menu	
Export	'Raw_Variables.py'	Python file containing a dictionary for each of the arrivals, service, capacity and service options used for each simulation. The dictionary keys are the [SimName].
Save	'Results_Tables.xlsx'	The four results tables each saved as a sheet. This will be overwritten on each save.

7 Glossary

Term	Description					
General						
Data	The selected input data.					
I	The number of individuals in the data.					
Dataframe	The dataset generated of the unique pathways within the data. This will be sorted by decreasing count then alphabetically decreasing.					
i	The number of entries in the dataframe which corresponds to the number of unique pathways in the data.					
Pathway	String of single character codes representing the chronological order of activities performed by an individual.					
Pathways Indexes	The number corresponding to the position of the pathway in the dataframe, starting at 0.					
Day	The time unit of a day.					
Named Day	Monday, Tuesday, Wednesday, Thursday, Friday, Saturday, Sunday. Day 0 will be Monday in the simulation.					
N	Number of activities.					
A_n	Number of arrivals at node n i.e. the number of times node n was the first activity in the pathway.					
P	Number of days in the period covered in the data.					
Floor	Formally the largest integer no larger than x [27] i.e. the integer.					
Ceiling	Formally the smallest integer no smaller than $x [27]$ i.e. the integer $+ 1$.					
Networks	Depiction of the Transition Matrix in the form of a Directed Graph.					
	Metrics					
Distance Metrics (code)	Methods used to calculate a numerical value to represent how different two strings are. The Python library Textdistance [21] is used to calculate all metrics except for the Modified Needleman-Wunsch.					
Levenshtein (Lev)	[32, 16]					
Damerau-Levenshtein (DLev)	[24, 9]					
Jaro (Jaro) Jaro-Winkler (JaroW)	[30, 13, 14]					
Needleman-Wunsch (NW)	[30, 15] [35, 17]					
Jaccard (Jac)	[29]					
Cosine (Cos)	[23, 8]					
Longest Common Subsequence (LCS)	[34]					
Modified Needleman-Wunsch (MNW)	Modification of the Needleman-Wunsch algorithm where the calculation considers rankings and groupings of the single character codes.					

Term	Description				
Clustering					
Transition Matrix	Matrix where position (i, j) represents the exact number of times activity i followed by activity j was performed.				
Adjusted Transition Matrix	Same as Transition Matrix but where any entries with values less than the Adjust Percentage are reduced to 0.				
Adjust Percentage	Percentage applied to the number of unique pathways e.g. 5% of 200 pathways = 10.				
Transition Probability Matrix	Matrix where position (i, j) represents the probability of activity i followed by activity j was performed. Calculated by each value in the Transition Matrix divided by the sum of its row.				
Difference Matrix	The total, mean and largest transitions are calculated by comparing the original transition matrix with the simulation transition matrix, not including the start and end activities.				
Distance Matrix	Matrix where position (i, j) represents the distance between string i and j .				
Propagated Values	As the clustering is performed for the dataframe (only unique pathways), when using the clustering results for the simulation one must consider 'propagating' the results to reflect the full data. For example, if pathway 'ABC' was assigned to cluster 0 and repeated 4 times in the original data, then the list of pathways for cluster 0 would contain ['ABC', 'ABC', 'ABC', 'ABC'].				
k-Medoids	Clustering method to separate the set of pathways into k clusters with one of the pathways as the centre point for each cluster. Points are assigned to the cluster with the smallest distance to the centre point (medoid). [31]				
Medoids	The pathway selected as the centre point for a cluster.				
k	The number of clusters.				
Max k	The maximum number of clusters to calculate.				
Initial Centroids	Initial starting centres for the k-medoids clustering.				
Silhouette Score	A metric to represent how 'good' the clustering is - used to suggest the number of clusters (k) to select. Describes how a point is to its asigned cluster centre compared to the centres of the other clusters [36]. Values range between -1 and 1 where a larger values indicates a better clustering. The Python library scikit-learn is used to calculate the silhouette score [19].				
Network Connections	Number of non-zero entries in the Transition Matrix				
Tolerance	Value to define the results to highlight i.e. if the number of network connections are within the selected tolerance of the number of network connections of the original data then results will be highlighted for that value of k.				

Term	Description				
Simulation					
Seed	As Ciw [3] uses random number generators, a seed is set to ensure that the same set of random numbers are used. This allows the simulation to be run multiples times and not have the results change due to the random numbers generated.				
Node	A node is an object where individuals are serviced. Each activity will have a corresponding node. Node count starts at 1.				
Dummy Node	A dummy node is used when the service at the node is unimportant but the node is required to structure the system.				
Class	A class is used to denote groups of individuals [4].				
Arrival Rate	The rate at which individuals will arrive into the system through the arrival node/s. An Exponential Distribution is use where the general arrival rate used is calculated by $\lambda = \frac{A_n}{P} \tag{1}$				
Exponential Distribution	The exponential distribution is used the sample the time between events which occur at an average rate λ [5, 26]. The exponential distribution is commonly used for the arrival rate.				
Deterministic Distribution	The deterministic distribution will always sample the same value [5].				
Warm Up	A warm up time is the time where the simulation will run without collecting results. This is to allow the simulation to 'fill up' before observation of the system. [7]				
W	For the iterative warm up, w is the number of times to simulate I individuals where only the final I individuals will record results e.g. with $w=3$ and $I=200$, the simulation will run for 600 individuals where only the final 200 (from id number 401 onwards) will be recorded.				

Plots

Bar Chart	A bar chart is used to represent categorical data [22, 11].
Directed Graph	A graph consisting of vertices and directed edges [25]. The directed graph represents the network of the activity interactions in the pathways. A vertex represents a node/activity and a directed edge represents the precedence of the vertex. The python library Graphviz [10] was used to create these Directed Graphs.
Histogram	A plot of the approximate distribution for numerical data [28]. Represents the frequency of occurrences. [11]
Line Chart	A series of data points joined together with a line [33, 11].
Violin Plot	Displays the distribution of the data highlighting the mean. [37, 11, 20]

References

- [1] E.F., Arruda, P. Harper, T. England, D. Gartner, E. Aspland, F.O, Ourique, T. Crosby. Resource optimization for cancer pathways with aggregate diagnostic demand: a perishable inventory approach *IMA Journal of Management Mathematics*, 2020.
- [2] Andrei Novikov. PyClustering: Data mining library. Journal of Open Source Software, 4(36):1230, April 2019.
- [3] The Ciw library developers Ciw: 2.0.1, 2019 https://github.com/CiwPython/Ciw
- [4] Ciw Class, The Ciw library developers Ciw: 2.0.1, 2019 https://ciw.readthedocs.io/en/latest/Tutorial-II/tutorial_vii.html
- [5] Ciw Distributions, The Ciw library developers Ciw: 2.0.1, 2019 https://ciw.readthedocs.io/en/latest/Guides/set_distributions.html
- [6] Ciw seed, The Ciw library developers Ciw: 2.0.1, 2019 https://ciw.readthedocs.io/en/latest/Guides/seed.html
- [7] Ciw Warm up, The Ciw library developers Ciw: 2.0.1, 2019 https://ciw.readthedocs.io/en/latest/Tutorial-I/tutorial_iv.html
- [8] P-N. Tan, M. Steinbach and V. Kumar. Introduction to Data Mining. Chapter 8, page 500, 2005.
- [9] F.J. Damerau. A technique for computer detection and correction of spelling errors. *Communications of the ACM*, Vol 7, Issue 3:171–176, 1964.
- [10] Graphviz, Python library, https://graphviz.readthedocs.io/en/stable/index.html
- [11] Hunter, J. D., Matplotlib: A 2D graphics environment, Computing in Science & Engineering, 9:3 90–95, 2007, https://matplotlib.org/
- [12] Matplotlib Navigation Toolbar, Matplotlib: A 2D graphics environment, https://matplotlib.org/3.1.1/users/navigation_toolbar.html
- [13] M.A. Jaro. Advances in Record-Linkage Methodology as Applied to Matching the 1985 Census of Tampa, Florida. Journal of the American Statistical Association, Volume 84, Issue 406, 1989.
- [14] M.A. Jaro. Probabilistic linkage of large public health data files. Statistics in Medicine, Volume 14, Issue 5-7, 1995.
- [15] W.E. Winkler. String Comparator Metrics and Enhanced Decision Rules in the Fellegi-Sunter Model of Record Linkage. Bureau of the Census, 1990.
- [16] V.I. Levenshtein. Binary Codes Capable Of Correcting Deletions, Insertions, and Reversals. Cybernetics and Control Theory, Vol 10, NO.8, 1966.
- [17] S.B. Needleman and C.D. Wunsch. A general method applicable to the search for similarities in the amino acid sequence of two proteins. *Journal of Molecular Biology*, Volume 48, Issue 3:443–453, 1970.
- [18] F. Pedregosa, G. Varoquaux, A. Gramfort, V. Michel, B. Thirion, O. Grisel, M. Blondel, P. Prettenhofer, R. Weiss, V. Dubourg, J. Vanderplas, A. Passos, D. Cournapeau, M. Brucher, M. Perrot, and E. Duchesnay. Scikit-learn: Machine learning in Python. Journal of Machine Learning Research, 12:2825–2830, 2011.
- [19] Silhouette Score, Scikit-Learn, https://scikit-learn.org/stable/modules/generated/sklearn.metrics.silhouette_score.html
- [20] Scikit-learn documentation for Violin Plots, Scikit-Learn, https://scikit-learn.org/stable/modules/density.html
- [21] Textdistance. Textdistance. Python package, 2017, https://pypi.org/project/textdistance/.
- [22] Bar Chart, Wikipedia, https://en.wikipedia.org/wiki/Bar_chart
- [23] Cosine Similarity, Wikipedia, https://en.wikipedia.org/wiki/Cosine_similarity

- [24] Damerau-Levenshtein Distance, Wikipedia, https://en.wikipedia.org/wiki/Damerau%E2%80%93Levenshtein_distance
- [25] Directed Graph, Wikipedia, https://en.wikipedia.org/wiki/Directed_graph
- [26] Exponential Distribution, Wikipedia, https://en.wikipedia.org/wiki/Exponential_distribution
- [27] Floor and Ceiling Functions, Wikipedia, https://en.wikipedia.org/wiki/Floor_and_ceiling_functions
- [28] Histogram, Wikipedia, https://en.wikipedia.org/wiki/Histogram
- [29] Jaccard Index, Wikipedia, https://en.wikipedia.org/wiki/Jaccard_index
- [30] Jaro Winkler Distance, Wikipedia, https://en.wikipedia.org/wiki/Jaro%E2%80%93Winkler_distance
- [31] k-Medoids Clustering, Wikipedia, https://en.wikipedia.org/wiki/K-medoids
- [32] Levenshtein Distance, Wikipedia, https://en.wikipedia.org/wiki/Levenshtein_distance
- [33] Line Chart, Wikipedia, https://en.wikipedia.org/wiki/Line_chart
- [34] Longest Common Subsequence, Wikipedia, https://en.wikipedia.org/wiki/Longest_common_subsequence_problem
- [35] Needleman Wunsch algorithm, Wikipedia, https://en.wikipedia.org/wiki/Needleman%E2%80%93Wunsch_algorithm
- [36] Silhouette Score, Wikipedia, https://en.wikipedia.org/wiki/Silhouette_(clustering)
- [37] Violin Plot, Wikipedia, https://en.wikipedia.org/wiki/Violin_plot